Page 1 of 7



PCT09

DATE: 02/06/2002 RAW SEQUENCE LISTING TIME: 14:15:13 PATENT APPLICATION: US/09/674,337A

Input Set : A:\55107sequence.txt

Output Set: N:\CRF3\02062002\I674337A.raw



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            Kyoko Higuchi
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            THE SAME
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                                                    140
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## DATE: 02/06/2002 RAW SEQUENCE LISTING PATENT APPLICATION: US/09/674,337A TIME: 14:15:13

Input Set : A:\55107sequence.txt

Output Set: N:\CRF3\02062002\1674337A.raw

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DATE: 02/06/2002

TIME: 14:15:13

RAW SEQUENCE LISTING PATENT APPLICATION: US/09/674,337A

Input Set : A:\55107sequence.txt

Output Set: N:\CRF3\02062002\1674337A.raw

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116	Ala	His	Tyr	Ser	Asp	Met	Leu	Ala	Ala	Phe	Asp	Lys	Pro	Leu	Asp	His
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119		100 105 110														
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122	Arq	Val	Ala	Phe	Ile	Gly	Ser	Gly	Pro	Leu	Pro	Phe	Ser	Ser	Phe	Val
123	3	130				-	135	-				140				
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133	Ory	210	niiu	mu	Olu	nop.	215	1114				220		Ļ	011	
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140	Ser	Val	Tle	Ile	Δla	Gln	T.vs	Ser		Asp	Val	His	λla		Glv	Leu
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DATE: 02/06/2002 RAW SEQUENCE LISTING TIME: 14:15:13 PATENT APPLICATION: US/09/674,337A

Input Set : A:\55107sequence.txt

Output Set: N:\CRF3\02062002\I674337A.raw

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163	cgcc	acct	gc c	ccgacaccat gtt				cgacaac tatgacctgt			gcgg	540							
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165	gacg	rtege	gg a	acctcgccgg (			gaget	cgcc	aag	aagtacgacg			ttgtcttcct ggccgcactc						
166	gtcg	gcat	.gg c	ccgccgagga			aggo	gaag	gtg	gatcg	gcgc	acct	cggc	gc a	caca	ıtggca	720		
167	gacg	igggo	gg c	ccto	gtcg	rt go	gcag	rcgca	cac	ggag	jcgc	gcgg	gtto	ct g	tacc	ccgatc	780		
168													tggccgtgtg ccatcccgac						
169													aggacgtgca tgccgatgga						
170										tacgcgcggg							960		
171										gacgtgaccc							1020		
172	ttto	rccaa	aca d	ccaa	ataa	rc ct	cttttqatcq			ttcgctgcga			gtgo	at d	cate	gatcca	1080		
173													atgcatttca agtcacgtgt						
174	tact	teta	atc c	aata	atαt	a co	catataatat			ttacacaca			ctto	ttgta	1200				
175	tgcttctatc caataatgta cgt							iaσta	cat	ataa	taa	tcac	1260						
176		tgtgtacaag tgaattttaa ttcacaagta catataatgg tcaccattga aaagatgttt agtgtgtgtt ttccaatata tgtttgtgta aggttcatca tctaataaaa tatgtttgga											1320						
177			aaa a					, - 5		,							1342		
	<210>					ia ac	•												
	<211>				_											•			
	<212>																		
	<213>				iorde	7 mile	ກນໄດຂ	re I											
	<400>					-uiii v	urge		- •										
185						λen	λen	Lys	Δen	Val	Δla	λla	Leu	Val	Glu	Lvs			
186	1	Ата	ніа	GIII	5	ASII	ASII	цуз	пор	10	1114		200		15				
187		mb r	C1 17	Tau	-	λla	λ1 =	Ile	Δla		T.e.ii	Pro	Ser	Leu		Pro			
188	116	1111	GIY	20	шта	AIG	пла	110	25	цуз	шец	110	001	30	001				
189	cor	Dro	N c n		λαη	λla	Leu	Phe		Glu	T.e.11	Val	Thr		Cvs	Val			
190	ser	PIO	35	val	АБР	мта	пец	40	T11T	OLU	БСС	vai	45	1114	0,15	, 41			
	Dmo	Dwo		Dro	17 - 1	λοη	37 - 1	Thr	Tvc	T (2))	Clv	Pro		Δla	Gln	Glu			
191	PIO	50	ser	PIO	vai	кър	55	1111	шуз	пси	GLY	60	Ora	mu	0111	014			
192	Mo+		C1.,	C1,,	T 011	T10		Leu	Cvc	Sor	Glu		Glu	Glv	T.VS	T.eu			
193		AIG	GIU	GTĀ	ьец	70	AIG	пеп	Cys	361	75	AIG	GIU	OLY	175	80			
194	65	7 l -	77.5 ~	(T)====	C		Mot	Leu	7. 7. 7.	אן א		7 cn	Nan	Dro	T. 🗕 11				
195	GIU	Ala	HIS	Tyr		Asp	Met	ьeu	Ald	90	Pne	ASP	ASII	FIO	95	мэр			
196	<b></b> .	_	<b>a</b> 1	<b>-</b> 1 -	85 Black	D	Ш	m	a		Шттт	т1 о	Nan	T 011		Two			
197	HIS	Leu	GIA		Pne	Pro	TÀL	Tyr		ASI	тАт	iie	ASII		261	цуѕ			
198	_		_	100	_	_		•	105	**- 1	3	3	TT -	110	Dwo	7.1 ~			
199	Leu							Arg							PIO	Ald			
200								120							<b>5</b> 1	**- 1			
201	Arg			Phe	Ile	Gly		Gly	Pro	Leu	Pro		ser	Ser	Pne	vai			
202		130					135					140	_	_	_	_			
203		Ala	Ala	Arg	His		Pro	Asp	Thr	Met			Asn	чуr	Asp				
204	145					150					155		_	<u>.</u> -	_	160			
205	Cys	Gly	Ala	Ala		Asp	Arg	Ala	Ser		Leu	Phe	Arg	Ala		Thr			
206					165					170					175	_			
207	Asp	Val	Gly	Ala	Arg	Met	Ser	Phe			Ala	Asp	Val		Asp	Leu			
208				180					185					190					
209	Ala	Ser	Glu	Leu	Ala	Lys	Tyr	Asp	Val	Val	Phe	Leu	Ala	Ala	Leu	Val			





DATE: 02/06/2002 RAW SEQUENCE LISTING TIME: 14:15:13 PATENT APPLICATION: US/09/674,337A

Input Set : A:\55107sequence.txt

Output Set: N:\CRF3\02062002\I674337A.raw

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205
            195
                                200
210
    Gly Met Ala Ala Glu Asp Lys Ala Lys Val Ile Ala His Leu Gly Ala
211
                                                220
212
                            215
    His Met Ala Asp Gly Ala Ala Leu Val Val Arg Ser Ala His Gly Ala
213
214
                        230
                                             235
    Arg Gly Phe Leu Tyr Pro Ile Val Asp Pro Gln Asp Ile Gly Arg Gly
215
                                        250
                    245
216
     Gly Phe Glu Val Leu Ala Val Cys His Pro Asp Asp Asp Val Val Asn
217
218
                                    265
                                                        270
     Ser Val Ile Ile Ala Gln Lys Ser Lys Glu Val His Ala Asp Gly Leu
219
                                 280
220
    Gly Ser Ala Arg Gly Ala Gly Arg Gln Tyr Ala Arg Gly Thr Val Pro
221
                             295
                                                 300
222
     Val Val Ser Pro Pro Cys Arg Phe Gly Glu Met Val Ala Asp Val Thr
223
                                             315
224
                         310
    Gln Asn His Lys Arg Asp Glu Phe Ala Asn Ala Glu Val Ala Phe
225
                     325
                                         330
226
228 <210> SEQ ID NO: 6
229 <211> LENGTH: 1314
230 <212> TYPE: DNA
231 <213> ORGANISM: Hordeum vulgare L.
233 <400> SEQUENCE: 6
    ctacttcact cacactagtg cccagaaaga aggctgcaat ggctgcccag aacaacaaca
                                                                            60
     aggatgtcgc tgccctggtg gagaagatca ccgggctcca cgccgccatc gccaagctgc
                                                                           120
                                                                           180
     cqtcqctcaq cccatccccq gacgtcgacg cgctcttcac cgagctggtc acggcgtgcg
236
                                                                           240
     ttcccccgag ccccgtggac gtgaccaagc tcggccccga ggcgcaggag atgcgggagg
237
     gcctcatccg cctctgctcc gaggccgagg ggaagctgga ggcgcactac tccgacatgc
                                                                           300
238
                                                                           360
239
     tegeogeett egacaacceg etggateace teggeatett eccetactae ageaactaea
     tcaacctcag caagctggag tacgagctcc tggcacgcta cgtccggcgg catcgcccgg
                                                                           420
240
                                                                           480
     cccgcgtcgc gttcatcggc tccggcccgc tgccgttcag ctcctttgtc ctggccgcgc
241
                                                                           540
     gccacctgcc cgacaccatg tttgacaact acgacctttg cggcgcggcc aacgatcgcg
242
                                                                           600
    ccaqcaagct cttccgcgcg gacacggacg tgggtgcccg catgtcgttc cacacggccg
243
     acgtcgcgga cctcgccagc gagctcgcca agtacgacgt cgtcttcctg gccgcgctcg
                                                                           660
244
     teggcatgge egecgaggae aaggecaagg tgategegea eeteggegea cacatggeag
                                                                           720
245
     acggggcggc cctcgtcgtg cgcagcgcac acggagcgcg cgggttcctg tacccgattg
                                                                           780
246
                                                                           840
     tcgaccccca ggacatcggc cgcggcgggt tcgaggtgct ggccgtgtgc caccccgacg
247
    acgacgtggt gaactccgtc atcatcgcac agaagtccaa ggaggtgcat gccgatggac
                                                                           900
248
    ttggcagcgc gcgtggtgcc ggtcgacagt acgcgcgcgg cacggtgccg gttgtcagcc
                                                                           960
249
                                                                          1020
250 ccccgtgcag gttcggtgag atggtggcgg atgtgaccca gaaccacaag agagacgagt
                                                                          1080
251 ttgccaacgc cgaagtggcc ttttgatcga tcgtcgccaa gggacaataa atgaacgtgg
                                                                          1140
    atgtggtagg gtaatttgcc tacctcgctg cttgatcgct tgcaatatgt gcacattttc
252
     ctactaccgc tgcttatgca tttcaagcca tgtgatgttg gtatccaata aagtatgtgt
                                                                          1200
253
                                                                          1260
     agggtttaca cgcaaatgtc tttacacctt gtacgtgtaa gtgttgacaa cgatgaattt
254
     1314
255
257 <210> SEQ ID NO: 7
258 <211> LENGTH: 329
259 <212> TYPE: PRT
260 <213> ORGANISM: Hordeum vulgare L.
262 <400> SEQUENCE: 7
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Use of n and/or Xaa has been detected in the Sequence Listing. harmatic laquence Listing to insure a corresponding Explanation is presented in the <220> to <223> fields of each sequence using n or Xaa.

.

DATE: 02/06/2002

TIME: 14:15:14

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/674,337A

Input Set : A:\55107sequence.txt

Output Set: N:\CRF3\02062002\1674337A.raw

L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date

 $L\!:\!541~M\!:\!341~W\!:$  (46) "n" or "Xaa" used, for SEQ ID#:14